SEQUENCE LISTING

(1) GENERAL INFORMATION:

    (i) APPLICANT: Mallet, Francois
              Oriol, Guy
              Mandrand, Bernard

   (ii) TITLE OF INVENTION: RNA AMPLIFICATION METHOD REQUIRING ONLY
        ONE MANIPULATION STEP

  (iii) NUMBER OF SEQUENCES: 13

   (iv) CORRESPONDENCE ADDRESS:
       (A) ADDRESSEE: OLIFF & BERRIDGE
       (B) STREET: P.O. Box 19928
       (C) CITY: Alexandria
       (D) STATE: Virginia
       (E) COUNTRY: USA
       (F) ZIP: 22320

    (v) COMPUTER READABLE FORM:
       (A) MEDIUM TYPE: Floppy disk
       (B) COMPUTER: IBM PC compatible
       (C) OPERATING SYSTEM: PC-DOS/MS-DOS
       (D) SOFTWARE: PatentIn Release #1.0, Version #1.25

   (vi) CURRENT APPLICATION DATA:
       (A) APPLICATION NUMBER: 08/825,617
       (B) FILING DATE: MARCH 31, 1997
       (C) CLASSIFICATION:

 (viii) ATTORNEY/AGENT INFORMATION:
       (A) NAME: Berridge, William P.
       (B) REGISTRATION NUMBER: 30,024
       (C) REFERENCE/DOCKET NUMBER: WPB 28662B

   (ix) TELECOMMUNICATION INFORMATION:
       (A) TELEPHONE: (703)836-6400
       (B) TELEFAX: (703)836-2787

(2) INFORMATION FOR SEQ ID NO:1:

    (i) SEQUENCE CHARACTERISTICS:
       (A) LENGTH: 21 base pairs
       (B) TYPE: nucleic acid
       (C) STRANDEDNESS: single
       (D) TOPOLOGY: linear

   (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

CCTATCTGTC CCCTCAGCTA C   21

(2) INFORMATION FOR SEQ ID NO:2:

    (i) SEQUENCE CHARACTERISTICS:
       (A) LENGTH: 18 base pairs
       (B) TYPE: nucleic acid
       (C) STRANDEDNESS: single
       (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

TCTATCAAAG CAACCCAC   18

(2) INFORMATION FOR SEQ ID NO:3:

        (i) SEQUENCE CHARACTERISTICS:
            (A) LENGTH: 19 base pairs
            (B) TYPE: nucleic acid
            (C) STRANDEDNESS: single
            (D) TOPOLOGY: linear

        (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

ATCTCTAGCA GTGGCGCCC   19

(2) INFORMATION FOR SEQ ID NO:4:

        (i) SEQUENCE CHARACTERISTICS:
            (A) LENGTH: 19 base pairs
            (B) TYPE: nucleic acid
            (C) STRANDEDNESS: single
            (D) TOPOLOGY: linear

        (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

TTCCTTCGGG CCTGTCGGG   19

(2) INFORMATION FOR SEQ ID NO:5:

        (i) SEQUENCE CHARACTERISTICS:
            (A) LENGTH: 22 base pairs
            (B) TYPE: nucleic acid
            (C) STRANDEDNESS: single
            (D) TOPOLOGY: linear

        (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

CACAAAATAG AGTGGTGGTT GC   22

(2) INFORMATION FOR SEQ ID NO:6:

        (i) SEQUENCE CHARACTERISTICS:
            (A) LENGTH: 22 base pairs
            (B) TYPE: nucleic acid
            (C) STRANDEDNESS: single
            (D) TOPOLOGY: linear

        (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

TCGTCGCTGT CTCCGCTTCT TC   22

(2) INFORMATION FOR SEQ ID NO:7:

        (i) SEQUENCE CHARACTERISTICS:
            (A) LENGTH: 20 base pairs
            (B) TYPE: nucleic acid

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            (C) STRANDEDNESS: single
            (D) TOPOLOGY: linear



       (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

CTTCCTGCCA TAGGAGATGC                                           20

(2) INFORMATION FOR SEQ ID NO:8:

       (i)·SEQUENCE CHARACTERISTICS:
            (A) LENGTH: 24 base pairs
            (B) TYPE: nucleic acid
            (C) STRANDEDNESS: single
            (D) TOPOLOGY: linear



       (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

CTCCATTTCT TGCTCTCCTC TGTC                                      24

(2) INFORMATION FOR SEQ ID NO:9:

       (i) SEQUENCE CHARACTERISTICS:
            (A) LENGTH: 23 base pairs
            (B) TYPE: nucleic acid
            (C) STRANDEDNESS: single
            (D) TOPOLOGY: linear



       (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

CCACACAACT ATTGCTATTA TTA                                       23

(2) INFORMATION FOR SEQ ID NO:10:

       (i) SEQUENCE CHARACTERISTICS:
            (A) LENGTH: 18 base pairs
            (B) TYPE: nucleic acid
            (C) STRANDEDNESS: single
            (D) TOPOLOGY: linear



       (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

AAGAAGCGGA GACAGCGA                                             18

(2) INFORMATION FOR SEQ ID NO:11:

       (i) SEQUENCE CHARACTERISTICS:
            (A) LENGTH: 20 base pairs
            (B) TYPE: nucleic acid
            (C) STRANDEDNESS: single
            (D) TOPOLOGY: linear



       (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

AGCAGAAGAC AGTGGCAATG                                           20

(2) INFORMATION FOR SEQ ID NO:12:
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(i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 23 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

AAAACCCACC TCCCAACCCC GAG   23

(2) INFORMATION FOR SEQ ID NO:13:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 22 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

GCAACCACCA CTCTATTTTG TG   22